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1: XP_057927. Reports ...[gi:20535708] This record was removed as a result of standard genome annotation processing. See the genome build documentation at <http://www.ncbi.nlm.nih.gov/genome/guide/build.html> for further information, or contact info@ncbi.nlm.nih.gov.

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LOCUS XP_057927 853 aa linear PRI 03-JAN-2003
 DEFINITION similar to KIAA1902 protein [Homo sapiens].
 ACCESSION XP_057927
 VERSION XP_057927.2 GI:20535708
 DBSOURCE REFSEQ: accession [XM_057927.3](#)
 KEYWORDS .
 SOURCE Homo sapiens (human)
 ORGANISM [Homo sapiens](#)
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 REFERENCE 1 (residues 1 to 853)
 AUTHORS International Human Genome Sequencing Consortium.
 TITLE The DNA sequence of Homo sapiens
 JOURNAL Unpublished (2003)
 COMMENT MODEL REFSEQ: This record is predicted by automated computational analysis. This record is derived from an annotated genomic sequence ([NT_005151](#)) using gene prediction method: BLAST, supported by mRNA and EST evidence.
 Also see:

[Documentation](#) of NCBI's Annotation Process

On May 13, 2002 this sequence version replaced [gi:16157374](#).

FEATURES
 Location/Qualifiers
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 /db_xref="taxon:9606"
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 /note="FH2; Formin Homology 2 Domain"
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 /note="FH2; Formin Homology 2 Domain"
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 /db_xref="GeneID:114793"

ORIGIN

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181 ivelekqlmq rnkeldvvre iykdantqvh tlrkmvkeke eaiqrqstle kkihelekqg
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361 tssptvvfns glaavkikkp iktkfrmpvf nwvalkpni ngtvfneidd eriledlnvd
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481 ihvfdlktlp vdfveclmrf lptenevkv lryererkpl enlsdedrfm mqfskierlm
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661 lldvkelqrg mdltkreytm hdhntllkef ilnnegklkk lqddakiaqd afddvvykfg
721 enpkktpsv ffpvfvrfrvk aykqaeene lrkkqeqalm ekllqealm eqqdpkpsph
781 kskrqqqeli aelrrrqvkd nrhvyegkdg alediidlr ngpyrradav rrsvrrrdd
841 qnlrsvngae itm

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